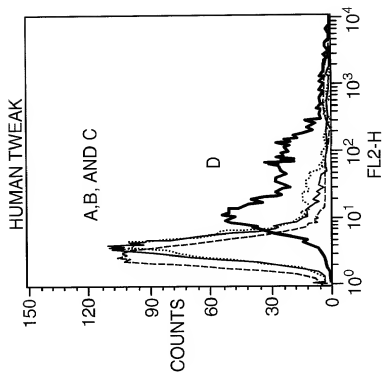
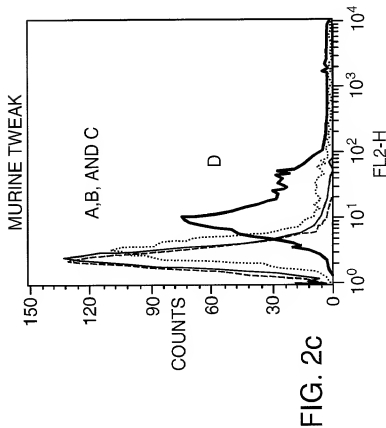
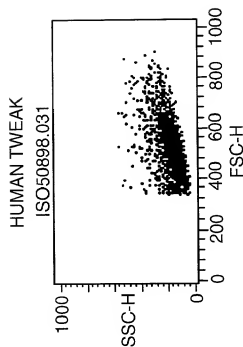
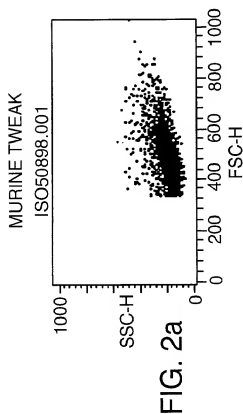


Human	1	MAARRSQRRRRGEPCGTALLVPLALGLGLALACLGLLLA	VVSLGSRASL
Mouse	1	-----	VLSLGLALACLGLLLVVSLGSWATL
Human	51	SAQEPAQEELVAEEDQDPSEINLPQTEESQDPAPFLNRLV	RRRSAPKGRK
Mouse	27	SAQEPSQEELTAEDRRPEPELNQTEESQDVVPFLEQLV	RRRSAPKGRK
Human	101	TRARRAIAAHYEVHPRPGQDGAQAGVDGTVSGWEEARIN	SSSPLRYNRQI
Mouse	77	ARPRRAIAAHYEVHPRPGQDGAQAGVDGTVSGWEETKIN	SSSPLRYDRQI
Human	151	GEFIVTRAGLYTYLYCQVHFDEGKAVYVKLDLLVDGVL	ALRCLEEPSATAA
Mouse	127	GEFTVIRAGLYTYLYCQVHFDEGKAVYVKLDLLVNGVL	ALRCLEEPSATAA
Human	201	SSLGPKQLRLCQVSGLLALRPGSSLRIRTL	PWAHLKAAPFLTYFGLFQVH
Mouse	177	SSPGPKQLRLCQVSGLLPLRPGSSLRIRTL	PWAHLKAAPFLTYFGLFQVH

FIG. 1



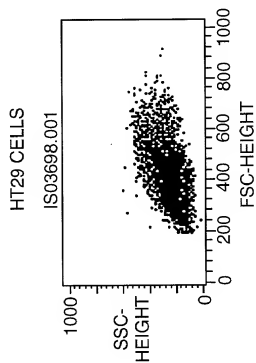


FIG. 3a

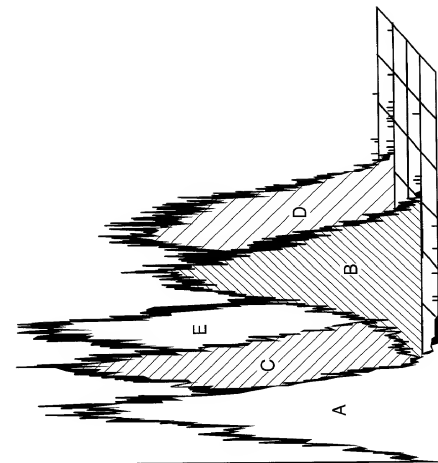


FIG. 3b

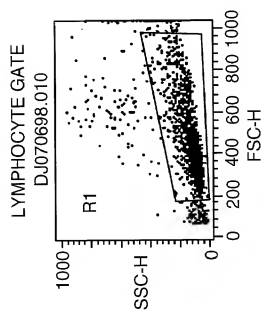


FIG. 4a

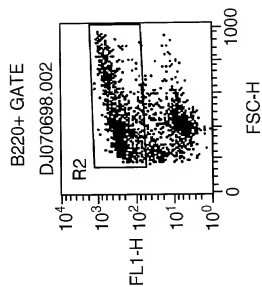


FIG. 4b

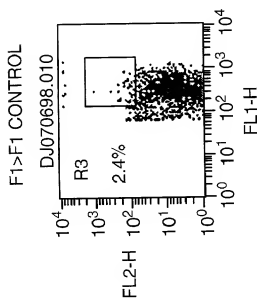


FIG. 4c

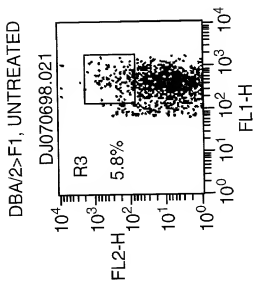


FIG. 4d

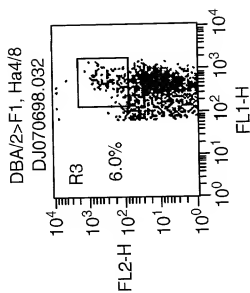


FIG. 4e

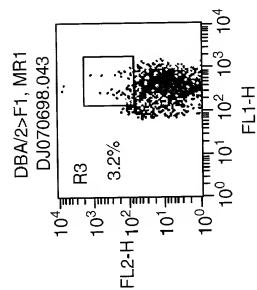


FIG. 4f

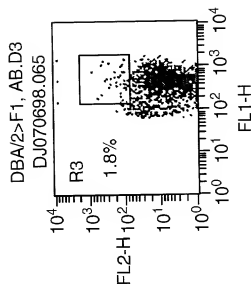


FIG. 4g